

10530789

=> d his all

(FILE 'HOME' ENTERED AT 15:55:35 ON 10 FEB 2010)

FILE 'REGISTRY' ENTERED AT 15:56:56 ON 10 FEB 2010

L1 STRUCTURE UPLOADED

L2 4 S L1

L3 58 S L1 FULL

L4 STRUCTURE UPLOADED

L5 33 S L4 FULL SUB=L3

FILE 'CAPLUS' ENTERED AT 15:59:10 ON 10 FEB 2010

L6 14 S L5

FILE 'STNGUIDE' ENTERED AT 15:59:50 ON 10 FEB 2010

FILE 'REGISTRY' ENTERED AT 16:04:29 ON 10 FEB 2010

L7 STRUCTURE UPLOADED

L8 34 S L7 FULL

FILE 'CAPLUS' ENTERED AT 16:04:56 ON 10 FEB 2010

L9 14 S L8

L10 0 S L9 NOT L6

FILE 'STNGUIDE' ENTERED AT 16:06:33 ON 10 FEB 2010

FILE 'REGISTRY' ENTERED AT 16:08:13 ON 10 FEB 2010

L11 1 S 681212-47-5/RN

L12 1 S 681212-50-0/RN

L13 1 S 681212-51-1/RN

L14 1 S 681212-52-2/RN

FILE 'STNGUIDE' ENTERED AT 16:09:18 ON 10 FEB 2010

FILE 'REGISTRY' ENTERED AT 16:11:49 ON 10 FEB 2010

L15 STRUCTURE UPLOADED

L16 1 S L15 FULL

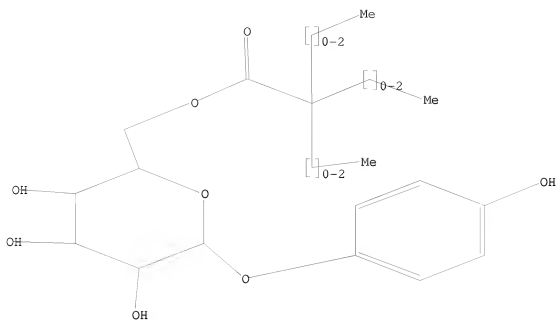
FILE 'CAPLUS' ENTERED AT 16:12:21 ON 10 FEB 2010

L17 2 S L16

=> d stat que

L15 STR

10530789



Structure attributes must be viewed using SIN Express query preparation.

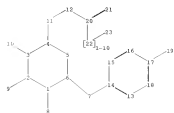
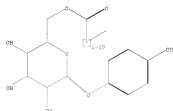
L16

1 SEA FILE=REGISTRY SSS FUL L15

L17

2 SEA FILE=CAPLUS ABB=ON PLU=ON L16

=> log h



```

chain nodes :
  7  8  9 10 11 12 19 20 21 22 23
ring nodes :
  1  2  3  4  5  6 13 14 15 16 17 18
chain bonds :
  1-8 2-9 3-10 4-11 6-7 7-14 11-12 12-20 17-19 20-21 20-22 22-23
ring bonds :
  1-2 1-6 2-3 3-4 4-5 5-6 13-14 13-18 14-15 15-16 16-17 17-18
exact/norm bonds :
  1-2 1-6 1-8 2-3 2-9 3-4 3-10 4-5 5-6 6-7 7-14 11-12 12-20 17-19 20-21
exact bonds :
  4-11 20-22 22-23
normalized bonds :
  13-14 13-18 14-15 15-16 16-17 17-18

```

```

Match level :
1:Atom 2:Atom 3:Atom 4:Atom 5:Atom 6:Atom 7:CLASS 8:CLASS 9:CLASS 10:CLASS 11:CLASS
12:CLASS 13:Atom 14:Atom 15:Atom 16:Atom 17:Atom 18:Atom 19:CLASS 20:CLASS 21:CLASS
22:CLASS 23:CLASS

```



```

chain nodes :
  7  8  9 10 11 12 19 20 21 22 23
ring nodes :
  1  2  3  4  5  6 13 14 15 16 17 18
chain bonds :
  1-8 2-9 3-10 4-11 6-7 7-14 11-12 12-20 17-19 20-21 20-22 22-23
ring bonds :
  1-2 1-6 2-3 3-4 4-5 5-6 13-14 13-18 14-15 15-16 16-17 17-18
exact/norm bonds :
  1-2 1-6 1-8 2-3 2-9 3-4 3-10 4-5 5-6 6-7 7-14 11-12 12-20 17-19 20-21
exact bonds :
  4-11 20-22 22-23
normalized bonds :
  13-14 13-18 14-15 15-16 16-17 17-18
  
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22:CLASS 23:CLASS
  
```

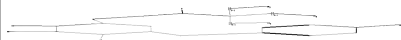


```

chain nodes :
  7  8  9 10 11 12 19 20 21 22 23
ring nodes :
  1  2  3 4  5  6 13 14 15 16 17 18
chain bonds :
  1-8 2-9 3-10 4-11 6-7 7-14 11-12 12-20 17-19 20-21 20-22 22-23
ring bonds :
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exact/norm bonds :
  1-2 1-6 1-8 2-3 2-9 3-4 3-10 4-5 5-6 6-7 7-14 11-12 12-20 17-19 20-21
exact bonds :
  4-11 20-22 22-23
normalized bonds :
  13-14 13-18 14-15 15-16 16-17 17-18
  
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1:Atom 2:Atom 3:Atom 4:Atom 5:Atom 6:Atom 7:CLASS 8:CLASS 9:CLASS 10:CLASS 11:CLASS
12:CLASS 13:Atom 14:Atom 15:Atom 16:Atom 17:Atom 18:Atom 19:CLASS 20:CLASS 21:CLASS
22:CLASS 23:CLASS
  
```



```

chain nodes :
  7  8  9 10 11 12 19 20 21 22 23 24 25 26 31 32
ring nodes :
  1  2  3  4  5  6 13 14 15 16 17 18
chain bonds :
  1-8 2-9 3-10 4-11 6-7 7-14 11-12 12-20 17-19 20-21 20-22 22-24 22-23 22-32 23-25
  24-26 31-32
ring bonds :
  1-2 1-6 2-3 3-4 4-5 5-6 13-14 13-18 14-15 15-16 16-17 17-18
exact/norm bonds :
  1-2 1-6 1-8 2-3 2-9 3-4 3-10 4-5 5-6 6-7 7-14 11-12 12-20 17-19 20-21
exact bonds :
  4-11 20-22 22-24 22-23 22-32 23-25 24-26 31-32
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  13-14 13-18 14-15 15-16 16-17 17-18

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Match level :
  1:Atom 2:Atom 3:Atom 4:Atom 5:Atom 6:Atom 7:CLASS 8:CLASS 9:CLASS 10:CLASS 11:CLASS
  12:CLASS 13:Atom 14:Atom 15:Atom 16:Atom 17:Atom 18:Atom 19:CLASS 20:CLASS 21:CLASS
  22:CLASS 23:CLASS 24:CLASS 25:CLASS 26:CLASS 31:CLASS 32:CLASS

```